

Figure 1

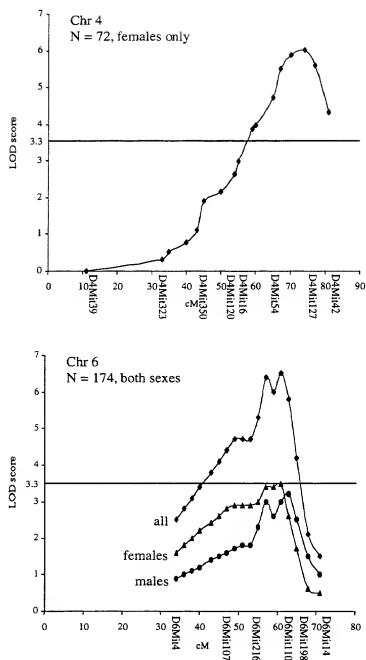


Figure 2

B-Isoform 1	351	AGAGCAGGAG	GAGCTTCTAC	AGAAGAATCA	GAACCTCCAA	GAAGCCCTGC
M-Isoform 1	351	AGAGCAGGAG	GAGCTTCTAC	AGAAGAATCA	GAACCTCCAA	GAAGCCCTGC
Isoform 7	73	-----	-----	-----	-----	-----
Isoform 8	73	-----	-----	-----	-----	-----
Isoform 9	73	-----	-----	-----	-----	-----
1st repeat						
B-Isoform 1	401	AAAGAGCTGC	AAACTCTTCA	GAGGAGTCCC	AGAGAGAACT	CAAGGGAAAAG
M-Isoform 1	401	AAAGAGCTGC	AAACTCTTCA	GAGGAGTCCC	AGAGAGAACT	CAAGGGAAAAG
Isoform 7	73	-----	-----	-AGGAGTCCC	AGAGAGAACT	CAAGGGAAAAG
Isoform 8	73	-----	-----	-----	-----	-----
Isoform 9	73	-----	-----	-----	-----	-----
1st repeat --> <---						
B-Isoform 1	451	ATAGACACCA	TCACCCGGAA	GCTGGACGAG	AAATCCAAAG	AGCAGGAGGA
M-Isoform 1	451	ATAGACACCA	TCACCCGGAA	GCTGGACGAG	AAATCCAAAG	AGCAGGAGGA
Isoform 7	102	ATAGACACCA	TCACCCGGAA	GCTGGACGAG	AAATCCAAAG	AGCAGGAGGA
Isoform 8	73	-----	-----	-----	-----	-----
Isoform 9	73	-----	-----	-----	-----	-----
2nd repeat						
B-Isoform 1	501	GCTTCTGCAG	ATGATTGAGA	ACCTCCAAGA	AGCCCTGCAG	AGAGCTGCAA
M-Isoform 1	501	GCTTCTGCAG	ATGATTGAGA	ACCTCCAAGA	AGCCCTGCAG	AGAGCTGCAA
Isoform 7	152	GCTTCTGCAG	ATGATTGAGA	ACCTCCAAGA	AGCCCTGCAG	AGAGCTGCAA
Isoform 8	73	-----	-----	-----	-----	-----
Isoform 9	73	-----	-----	-----	-----	-----
2nd repeat						
B-Isoform 1	551	ACTCTTCAGA	GGAGTCCCAG	AGAGAACTCA	AGGGAAAGAT	AGACACCCCTC
M-Isoform 1	551	ACTCTTCAGA	GGAGTCCCAG	AGAGAACTCA	AGGGAAAGAT	AGACACCCCTC
Isoform 7	202	ACTCTTCAGA	GGAGTCCCAG	AGAGAACTCA	AGGGAAAGAT	AGACACCCCTC
Isoform 8	73	-----A	GGAGTCCCAG	AGAGAACTCA	AGGGAAAGAT	AGACACCCCTC
Isoform 9	73	-----	-----	-----	-----	-----
2nd ----> <---- 3rd repeat						
B-Isoform 1	601	ACCTTGAAGC	TGAACGAGAA	ATCCAAAGAG	CAGGAGGAGC	TTCTACAGAA
M-Isoform 1	601	ACCTTGAAGC	TGAACGAGAA	ATCCAAAGAG	CAGGAGGAGC	TTCTACAGAA
Isoform 7	252	ACCTTGAAGC	TGAACGAGAA	ATCCAAAGAG	CAGGAGGAGC	TTCTACAGAA
Isoform 8	114	ACCTTGAAGC	TGAACGAGAA	ATCCAAAGAG	CAGGAGGAGC	TTCTACAGAA
Isoform 9	73	-----	-----	-----	-----	-----
3rd repeat						
B-Isoform 1	651	GAATCAGAAC	CTCCAAGAG	CCCTGCAAG	AGCTGCAAA	TTTTCAGGTC
M-Isoform 1	651	GAATCAGAAC	CTCCAAGAG	CCCTGCAAG	AGCTGCAAA	TTTTCAGGTC
Isoform 7	302	GAATCAGAAC	CTCCAAGAG	CCCTGCAAG	AGCTGCAAA	TTTTCAGGTC
Isoform 8	164	GAATCAGAAC	CTCCAAGAG	CCCTGCAAG	AGCTGCAAA	TTTTCAGGTC
Isoform 9	73	-----	-----	-----	-----	-----GTC
3rd repeat ----->						

Figure 3B

B-Isoform	1	701	CTTGTCCACA	AGACTGGCTC	TGGCATAAAG	AAAACGTGTTA	CCTCTTCCAT
M-Isoform	1	701	CTTGTCCACA	AGACTGGCTC	TGGCATAAAG	AAAACGTGTTA	CCTCTTCCAT
Isoform 7		352	CTTGTCCACA	AGACTGGCTC	TGGCATAAAG	AAAACGTGTTA	CCTCTTCCAT
Isoform 8		214	CTTGTCCACA	AGACTGGCTT	TGGCATAAAG	AAAACGTGTTA	CCTCTTCCAT
Isoform 9		75	CTTGTCCACA	AGACTGGCTC	TGGCATAAAG	AAAACGTGTTA	CCTCTTCCAT

B-Isoform	1	751	GGGCCCCCTTA	GCTGGGAAAA	AAACCGGCAG	ACCTGCCAAT	CITTTGGSTGG
M-Isoform	1	751	GGGCCCCCTTA	GCTGGGAAAA	AAACCGGCAG	ACCTGCCAAT	CITTTGGSTGG
Isoform 7		402	GGGCCCCCTTG	GCTGGGAAAA	AAACCGGCAG	ACCTGCCAAT	CITTTGGSTGG
Isoform 8		264	GGGCCCCCTTA	GCTGGGAAAA	AAACCGGCAG	ACCTGCCAAT	CITTTGGSTGG
Isoform 9		125	GGGCCCCCTTA	GCTGGGAAAA	AAACCGGCAG	ACCTGCCAAT	CITTTGGSTGG

B-Isoform	1	801	CCAGTTACTA	CAAATTAATG	GTGCAGATGA	TCTGACATTC	ATCTTACAAG
M-Isoform	1	801	CCAGTTACTA	CAAATTAATG	GTGCAGATGA	TCTGACATTC	ATCTTACAAG
Isoform 7		452	CCAGTTACTA	CAAATTAATG	GTGCAGATGA	TCTGACATTC	ATCTTACAAG
Isoform 8		314	CCAGTTACTA	CAAATTAATG	GTGCAGATGA	TCTGACATTC	ATCTTACAAG
Isoform 9		175	CCAGTTACTA	CAAATTAATG	GTGCAGATGA	TCTGACATTC	ATCTTACAAG

B-Isoform	1	851	CAATTTCCCA	TACCACCTCC	CCGTTCTGGA	TTGGATTGCA	TGGGAAGAAG
M-Isoform	1	851	CAATTTCCCA	TACCACCTCC	CCATTCCTGGA	TTGGATTGCA	TGGGAAGAAG
Isoform 7		502	CAATTTCCCA	TACCACCTCC	CCATTCCTGGA	TTGGATTGCA	TGGGAAGAAG
Isoform 8		364	CAATTTCCCA	TACCACCTCC	CCATTCCTGGA	TTGGATTGCA	TGGGAAGAAG
Isoform 9		225	CAATTTCCCA	TACCACCTCC	CCATTCCTGGA	TTGGATTGCA	TGGGAAGAAG

B-Isoform	1	901	CCTGGCCAAC	CATGGCTATG	GGAGAATGGA	ACTCCTTTGA	ATTTTCAATT
M-Isoform	1	901	CCTGGCCAAC	CATGGCTATG	GGAGAATGGA	ACTCCTTTGA	ATTTTCAATT
Isoform 7		552	CCTGGCCAAC	CATGGCTATG	GGAGAATGGA	ACTCCTTTGA	ATTTTCAATT
Isoform 8		414	CCTGGCCAAC	CATGGCTATG	GGAGAATGGA	ACTCCTTTGA	ATTTTCAATT
Isoform 9		275	CCTGGCCAAC	CATGGCTATG	GGAGAATGGA	ACTCCTTTGA	ATTTTCAATT

B-Isoform	1	951	CTTTAAGACC	AGGGGGCGTT	CTTTACAGCT	ATATTCAATCA	GGCAACTGTG
M-Isoform	1	951	CTTTAAGACC	AGGGGGCGTT	CTTTACAGCT	ATATTCAATCA	GGCAACTGTG
Isoform 7		602	CTTTAAGACC	AGGGGGCGTT	CTTTACAGCT	ATATTCAATCA	GGCAACTGTG
Isoform 8		464	CTTTAAGACC	AGGGGGCGTT	CTTTACAGCT	ATATTCAATCA	GGCAACTGTG
Isoform 9		325	CTTTAAGACC	AGGGGGCGTT	CTTTACAGCT	ATATTCAATCA	GGCAACTGTG

B-Isoform1	1001	CATACCTTCA	AGACGGAGCT	GTGTTGCGTG	AAAACGTGAT	TCTAATTGCA
M-Isoform1	1001	CATACCTTCA	AGACGGAGCT	GTGTTGCGTG	AAAACGTGAT	TCTAATTGCA
Isoform 7	652	CATACCTTCA	AGACGGAGCT	GTGTTGCGTG	AAAACGTGAT	TCTAATTGCA
Isoform 8	514	CATACCTTCA	AGACGGAGCT	GTGTTGCGTG	AAAACGTGAT	TCTAATTGCA
Isoform 9	375	CATACCTTCA	AGACGGAGCT	GTGTTGCGTG	AAAACGTGAT	TCTAATTGCA

B-Isoform1	1051	TTCAGCATAT	GTCAGAAGAA	GACAAATCAT	TTGCAAAATT	AG-----
M-Isoform1	1051	TTCAGCATAT	GTCAGAAGAA	GACAAATCAT	TTGCAAAATT	AG-----
Isoform 7	702	TTCAGCATAT	GTCAGAAGAA	GACAAATCAT	TTGCAAAATT	AG-----
Isoform 8	564	TTCAGCATAT	GTCAGAAGAA	GACAAATCAT	TTGCAAAATT	AG-----
Isoform 9	425	TTCAGCATAT	GTCAGAAGAA	GACAAATCAT	TTGCAAAATT	AG-----

Figure 3C

Isoform 1															
atg	act	ttt	gat	gac	aag	atg	aag	cct	gag	aat	gac	gag	cct	gat	cag
Met	Thr	Phe	Asp	Asp	Lys	Met	Lys	Pro	Ala	Asn	Asp	Glu	Pro	Asp	Gln
			5					10					15		
aag	tca	tgt	ggc	aag	aag	cct	aaa	ggt	ctg	cat	tgg	ctt	tct	tcc	cca
Lys	Ser	Cys	Gly	Lys	Lys	Pro	Lys	Gly	Leu	His	Leu	Leu	Ser	Ser	Pro
			20					25					30		
tgg	tgg	ttc	act	gct	gct	atg	act	ctg	gtc	atc	ctc	tgc	ctg	gtg	tgg
Trp	Trp	Phe	Pro	Ala	Ala	Met	Thr	Leu	Val	Ile	Leu	Cys	Leu	Val	Leu
			35				40				45				
tca	gtg	acc	ctt	att	gta	cag	tgg	aca	caa	tta	cgc	cag	gta	tct	gac
Ser	Val	Thr	Leu	Ile	Val	Gln	Trp	Thr	Gln	Leu	Arg	Gln	Val	Ser	Asp
	50					55					60				
ctc	tta	aaa	caa	tac	caa	gag	aac	ctt	act	cag	cag	gat	cgt	atc	ctg
Leu	Leu	Lys	Gln	Tyr	Gln	Ala	Asn	Leu	Thr	Gln	Gln	Asp	Arg	Ile	Leu
65				70						75				80	
gaa	ggg	cag	atg	tta	gcc	cag	cag	aag	gca	gaa	aac	act	tca	cag	gaa
Glu	Gly	Gln	Met	Leu	Ala	Gln	Gln	Lys	Ala	Glu	Asn	Thr	Ser	Gln	Glu
				85					90				95		
tca	aag	aag	gaa	ctg	aaa	gga	aag	ata	gac	acc	ctc	acc	cag	aag	ctg
Ser	Lys	Lys	Glu	Leu	Lys	Gly	Lys	Ile	Asp	Thr	Leu	Thr	Gln	Lys	Leu
			100					105					110		
aac	gag	aaa	tcc	aaa	gag	cag	gag	gag	ctt	cta	cag	aag	aat	cag	aac
Asn	Glu	Lys	Ser	Lys	Glu	Gln	Glu	Glu	Leu	Leu	Gln	Lys	Asn	Gln	Asn
		115				120					125				
ctc	caa	gaa	gcc	ctg	caa	aga	gct	gca	aac	tct	tca	gag	gag	tcc	cag
Leu	Gln	Glu	Ala	Leu	Gln	Arg	Ala	Ala	Asn	Ser	Ser	Glu	Glu	Ser	Gln
	130					135					140				
aga	gaa	ctc	aag	gga	aag	ata	gac	acc	atc	acc	cgg	aag	ctg	gac	gag
Arg	Glu	Leu	Lys	Gly	Lys	Ile	Asp	Thr	Ile	Thr	Arg	Lys	Leu	Asp	Glu
	145				150					155				160	
aaa	tcc	aaa	gag	cag	gag	gag	ctt	ctg	cag	atg	att	cag	aac	ctc	caa
Lys	Ser	Lys	Glu	Gln	Glu	Glu	Leu	Leu	Gln	Met	Ile	Gln	Asn	Leu	Gln
			165					170					175		
gaa	gcc	ctg	cag	aga	gct	gca	aac	tct	tca	gag	gag	tcc	cag	aga	gaa
Glu	Ala	Leu	Gln	Arg	Ala	Ala	Asn	Ser	Ser	Glu	Glu	Ser	Gln	Arg	Glu
			180					185					190		
ctc	aag	gga	aag	ata	gac	acc	ctc	acc	tgg	aag	ctg	aac	gag	aaa	tcc
Leu	Lys	Gly	Lys	Ile	Asp	Thr	Leu	Thr	Leu	Lys	Leu	Asn	Glu	Lys	Ser
			195				200					205			
aaa	gag	cag	gag	gag	ctt	cta	cag	aag	aat	cag	aac	ctc	caa	gaa	gcc
Lys	Glu	Gln	Glu	Glu	Leu	Gln	Lys	Asn	Gln	Asn	Leu	Gln	Glu	Glu	Ala
	210				215						220				

Figure 4A

[illegible]

Figure 4B

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Isoform 2

atg act ttc gat gac aag atg aag cct ggc aat gac gag cct gat cag	48
Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Gln	
1 5 10 15	
aag tca tgt ggc aag aag cct aaa ggt ctg cat ttg ctt tct tcc cca	86
Lys Ser Cys Gly Lys Lys Pro Lys Gly Leu His Leu Leu Ser Ser Pro	
20 25 30	
tgg tgg ttc cot gct gct atg act ctg gtc atc ctc tgc ctg gtg ttg	144
Trp Trp Phe Pro Ala Ala Met Thr Leu Val Ile Leu Cys Leu Val Leu	
35 40 45	
tca gtg acc ctt att gta cag tgg aca caa tgatcgtatc ctggaagggc	184
Ser Val Thr Leu Ile Val Gln Trp Thr Gln	
50 55	
agatgttagc ccagcagaag gcagaaaaca cttcacagga atcaaagaag gaactgaaag	254
gaaaagataga caccctcacc cagaagctga acgagaaatc caaagagcag gaggagcttc	314
tacagaagaa tcagaacctc caagaagccc tgcaaagagc tgcaaaactct tcagaggagt	374
cccagagaga actcaaggga aagatagaca ccatcaccgc gaagctggac gagaaatcca	434
aagagcagga ggagcttctg cagatgattc agaacctcca agaagccctg cagagagctg	494
caaaactcttc agaggagtc cagagagAAC tcaagggaaa gatagacacc ctcaccttga	554
agctgaacga gaaatccaaa gagcaggagg agcttctaca gaagaatcag aacctccaag	614
aagcccttga aagagctgca aacttttcag gtccttgctc acaagactgg ctctggcata	674
aagaaaaactg ttacctcttc cgtgggcccct ttactgggaa aaaagccggc agacctgcca	734
atctttgggt ggcagttact acaaatataa ggcagatg	794

Figure 5

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Isoform 3

atg act ttt gat gac aag atg aag cct gcg aat gac gag cct gat cag	48
Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Gln	
1 5 10	
aag tca tgt ggc aag aag cct aaa ggt ctg cat ttg ctt tct tcc cca	46
Lys Ser Cys Gly Lys Lys Pro Lys Gly Leu His Leu Leu Ser Ser Pro	
20 25 30	
tgg tgg ttc cct gct gct atg act ctg gtc atc ctc tgc ctg gtg ttg	44
Trp Trp Phe Pro Ala Ala Met Thr Leu Val Ile Leu Cys Leu Val Leu	
35 40 45	
tca gtg acc ctt att gta cag tgg aca caa tta cgc cag gta tct gac	42
Ser Val Thr Leu Ile Val Gln Trp Thr Gln Leu Arg Gln Val Ser Asp	
50 55 60	
ctc tta aaa caa tac caa gcg aac ctt act cag cag gat cgt atc ctg	24
Leu Leu Lys Gln Tyr Gln Ala Asn Leu Thr Gln Gln Asp Arg Ile Leu	
65 70 80	
gaa ggg cag atg tta gcc cag cag aag gca gaa aac act tca ccg caa	40
Glu Gly Gln Met Leu Ala Gln Gln Lys Ala Glu Asn Thr Ser Pro Gln	
85 90 95	
tca aag aag gaa ctg aaa gga aag ata gac acc ctc acc cag aag ctg	336
Ser Lys Lys Glu Leu Lys Gly Lys Ile Asp Thr Leu Thr Gln Lys Leu	
100 105 110	
aac gag aaa tcc aaa gag cag gag gag ctt cta cag aag aat cag aac	384
Asn Glu Lys Ser Lys Glu Gln Glu Glu Leu Leu Gln Lys Asn Gln Asn	
115 120 125	
ctc caa gaa gcc ctg caa aga gct gca aac tct tca gag gag tcc cag	432
Leu Gln Glu Ala Leu Gln Arg Ala Ala Asn Ser Ser Glu Glu Ser Gln	
130 135 140	
aga gaa ctc aag gga aag ata gac acc ctc acc ttg aag ctg aac gag	480
Arg Glu Leu Lys Gly Lys Ile Asp Thr Leu Thr Leu Lys Leu Asn Glu	
145 150 155 160	
aaa tcc aaa gag cag	495
Lys Ser Lys Glu Gln	
165	

Figure 6

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Isoform 4																	
atg	act	ttt	gat	gac	aag	atg	aag	cct	gcg	aat	gac	gag	cct	gat	cag		72
Met	Thr	Phe	Asp	Asp	Lys	Met	Lys	Pro	Ala	Asn	Asp	Glu	Pro	Asp	Gln		
					5				10					15			
aag	tca	tgt	ggc	aag	aag	cct	aaa	ggt	ctg	cat	ttg	ctt	tct	tcc	cca		86
Lys	Ser	Cys	Gly	Lys	Lys	Pro	Lys	Gly	Leu	His	Leu	Leu	Ser	Ser	Pro		
			20					25					30				
tgg	tgg	ttc	cct	gct	gct	atg	act	ctg	gtc	atc	ctc	tgc	ctg	gtg	ttg		144
Trp	Trp		Pro	Ala	Ala	Met	Thr	Leu	Val	Ile	Leu	Cys	Leu	Val	Leu		
		35					40				45						
tca	gtg	acc	ctt	att	gta	cag	tgg	aca	caa	tta	cgc	cag	gta	tct	gac		192
Ser	Val	Thr	Leu	Ile	Val	Gln	Trp	Thr	Gln	Leu	Arg	Gln	Val	Ser	Asp		
	50					55					60						
ctc	tta	aaa	caa	tac	caa	gcg	aac	ctt	act	cag	cag	gat	cgt	atc	ctg		240
Leu	Leu	Lys	Gln	Tyr	Gln	Ala	Asn	Leu	Thr	Gln	Gln	Asp	Arg	Ile	Leu		
	65				70					75				80			
gaa	ggg	cag	atg	tta	gcc	cag	cag	aag	gca	gaa	aac	act	tca	cag	gaa		288
Glu	Gly	Gln	Met	Leu	Ala	Gln	Gln	Lys	Ala	Glu	Asn	Thr	Ser	Gln	Glu		
			85						90					95			
tta	aag	aag	gaa	ctg	aaa	gga	aag	ata	gac	acc	ctc	acc	cag	aag	ctg		336
Ser	Lys	Lys	Glu	Leu	Lys	Gly	Lys	Ile	Asp	Thr	Leu	Thr	Gln	Lys	Leu		
			100					105					110				
aac	gag	aaa	tcc	aaa	gag	cag	gag	gag	ctt	cta	cag	aag	aat	cag	aac		364
Asn	Glu	Lys	Ser	Lys	Glu	Gln	Glu	Glu	Leu	Leu	Gln	Lys	Asn	Gln	Asn		
		115				120					125						
ctc	caa	gaa	gcc	ctg	caa	aga	gct	gca	aac	ttt	tca	ggt	cct	tgt	cca		432
Leu	Gln	Glu	Ala	Leu	Gln	Arg	Ala	Ala	Asn	Phe	Ser	Gly	Pro	Cys	Pro		
		130				135					140						
caa	gac	tgg	ctc	tgg	cat	aaa	gaa	aac	tgt	tac	ctc	ttc	cat	ggg	ccc		480
Gln	Asp	Trp	Leu	Trp	His	Lys	Glu	Asn	Cys	Tyr	Leu	Phe	His	Gly	Pro		
	145				150					155				160			
ttt	agc	tgg	gaa	aaa	aac	cgg	cag	acc	tgc	caa	tct	ttg	ggt	ggc	cag		528
Phe	Ser	Trp	Glu	Lys	Asn	Arg	Gln	Thr	Cys	Gln	Ser	Leu	Gly	Gly	Gln		
			165						170					175			
tta	cta	caa	att	aat	ggt	gca	gat	gat	ctg	aca	ttc	atc	tta	caa	gca		576
Leu	Leu	Gln	Ile	Asn	Gly	Ala	Asp	Asp	Leu	Thr	Phe	Ile	Leu	Gln	Ala		
			180					185					190				
att	tcc	cat	acc	acc	tcc	ccg	ttc	tgg	att	gga	ttg	cat	cgg	aag			624
Ile	Ser	His	Thr	Thr	Ser	Pro	Phe	Trp	Ile	Gly	Leu	His	Arg	Lys			
			195				200					205					

Figure 7

Isoform 5

atg act ttt gat gac aag atg aag cct gcg aat gac gag cct gat gag	48
Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Glu	
1 5 10 15	
aag tca tgt ggc aag aag cct aaa ggt ctg cat ttg ctt tct tcc cca	96
Lys Ser Cys Gly Lys Lys Pro Lys Gly Leu His Leu Leu Ser Ser Pro	
20 25 30	
egg tgg ttc cct gct gct atg act ctg gtc atc ctc tgc ctg gtg ttg	144
Trp Trp Phe Pro Ala Ala Met Thr Leu Val Ile Leu Cys Leu Val Leu	
35 40 45	
tca gtg acc ctt att gta cag tgg aca caa tgatcgratc ctggaagggc	192
Ser Val Thr Leu Ile Val Gln Trp Thr Gln	
50 55	
agatgttagc ccagcagaag gcagaaaaca cttcacagga atcaaagaag gaactgaaag	254
gaaagataga caccctcacc cagaagctga acgactccaa agagcaggag gagctacacc	314
ccccccgaac ctccaagaag ccttgcaaag agctgcaaac tcttcaggtc cttgtccaca	374
agactggctc tggcataaag aaaactgtta cctcttccat gggcccttta gctgggaaaa	434
aaaccggcag acctgccaat ctttgggtgg gcagttacta caaattaatg gtgcagatga	494
tctgacattc atcttacaag caatttccca taccacctcc ctttcttgga ttggattgca	554
tgggaagaag cctggcaacc atgggtatgg gagaatggac ttctttgaat ttttaatttt	614
aagacagggc gtttttacag tttttcataa ggactttgtg tacttagagg gctggggttcg	674
ctgaaatgat tctattggtt agcatgtaga aaaaaatt	

Figure 8

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Isoform 6

atg act ttt gat gac aag atg aag cct gcg aat gac gag cct gat cag	48
Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Gln	
1 5 10 15	
aag tca tgt ggc aag aag cct aaa ggt ctg cat ttg ctt tct tcc oca	96
Lys Ser Cys Gly Lys Lys Pro Lys Gly Leu His Leu Leu Ser Ser Pro	
20 25 30	
tgg tgg ttc cct gct gct atg act ctg gtc atc ctc tgc ctg gtg ttg	144
Trp Trp Phe Pro Ala Ala Met Thr Leu Val Ile Leu Cys Leu Val Leu	
35 40 45	
tca gtg acc ctt att gta cag tgg aca caa taggagtcgc agagagaact	192
Ser Val Thr Leu Ile Val Gln Trp Thr Gln	
50 55	
caagggaag atagacaccc tcacottgaa gotgaacgag aaatocaaag agcaggagga	240
gottctacag aagaatcaga acctccaaga agccctgcaa agagctgcaa acttttcagg	288
tccttgctca caagactggc tctggcataa agaaaaactgt tacctcttcc atgggccctt	336
tagctgggaa aaaaaccggc agacctgcca atctttgggt ggccagttac taaaaattaa	384
tggcgagat gatctgacat tcactttaca agcaatttcc cataccacct ccccgttctg	432
gattggattg catcggaaga agccctggcca accatggcta tgggagaatg gaactccttt	480
gaattttcaa ttctttaaga ccaggggcgt ttctttacag ctatattcat caggcaactg	528
tgcatacctt caagacggac tgtgttcgct gaaaaactgca ttctaattgc attcagcata	576
tgtcaaaaga agacaaatca ttgcaaat ttgtgaatct aaagaat	624

Figure 9

Isoform 7

atg act ttt gat gac aag atg aag cct ggc aat gac gag cct gat cag
 Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Gln
 1 5 10 15

aag tca tgt ggc aag aag cct aaa gag gag tcc cag aga gaa ctc aag
 Lys Ser Cys Gly Lys Lys Pro Lys Glu Ser Gln Arg Glu Leu Lys
 20 25 30

gga aag ata gac acc atc acc cgg aag ctg gac gag aaa tcc aaa gag
 Gly Lys Ile Asp Thr Ile Thr Arg Lys Leu Asp Glu Lys Ser Lys Glu
 35 40 45

cag gag gag ctt ctg cag atg att cag aac ctc caa gaa gcc ctg cag
 Gln Gln Glu Leu Leu Gln Met Ile Gln Asn Leu Gln Glu Ala Leu Gln
 50 55 60

aga gct gca aac tct tca gag gag tcc cag aga gaa ctc aag gga aag
 Arg Ala Ala Asn Ser Ser Glu Glu Ser Gln Arg Glu Leu Lys Gly Lys
 65 70 75 80

ata gac acc ctc acc ttg aag ctg aac gag aaa tcc aaa gag cag gag
 Ile Asp Thr Leu Thr Leu Lys Leu Asn Glu Lys Ser Lys Glu Gln Glu
 85 90 95

gag ctt cta cag aag aat cag aac ctc caa gaa gcc ctg caa aga gct
 Glu Leu Leu Gln Lys Asn Gln Asn Leu Gln Glu Ala Leu Gln Arg Ala
 100 105 110

gca aac ttt tca ggt cct tgt cca caa gac tgg ctc tgg cat aaa gaa
 Ala Asn Phe Ser Ser Gly Pro Cys Pro Gln Asp Trp Leu Trp His Lys Glu
 115 120 125

aac tgt tac ctc ttc cat ggg ccc ttt ggc tgg gaa aaa aac cgg cag
 Asn Cys Tyr Leu Phe His Gly Pro Phe Gly Trp Glu Lys Asn Arg Gln
 130 135 140

acc tgc caa tct ttg ggt ggc cag tta cta caa att aat ggt gca gat
 Thr Cys Gln Ser Leu Gly Gly Gln Leu Leu Gln Ile Asn Gly Ala Asp
 145 150 155 160

gat ctg aca ttc atc tta caa gca att tcc cat acc acc tcc cca ttc
 Asp Leu Thr Phe Ile Leu Gln Ala Ile Ser His Thr Thr Ser Pro Phe
 165 170 175

tgg att gga ttg cat cgg aag aag cct ggc caa cca tgg cta tgg gag
 Trp Ile Gly Leu His Arg Lys Lys Pro Gly Gln Pro Trp Leu Trp Glu
 180 185 190

aat gga act cct ttg aat ttt caa ttc ttt aag acc agg ggc gtt tct
 Asn Gly Thr Pro Leu Asn Phe Gln Phe Phe Lys Thr Arg Gly Val Ser
 195 200 205

tta cag cta tat tca tca agc aac tgt gca tac ctt caa gac gga gct
 Leu Gln Leu Tyr Ser Ser Ser Asn Cys Ala Tyr Leu Gln Asp Gly Ala
 210 215 220

gtg ttc gct gaa aac tgc att cta att gca ttc agc ata tgt cag aag
 Val Phe Ala Glu Asn Cys Ile Leu Ile Ala Phe Ser Ile Cys Gln Lys
 225 230 235 240

aag aca aat cat ttg caa att tag
 Lys Thr Asn His Leu Gln Ile
 245

Figure 10

tgt acc ttt gat gaa aag atg aag cct gcg aat gac gag cct gat cag	48
Met Thr Phe Asp Asp Lys Met Lys Pro Ala Asn Asp Glu Pro Asp Gln	
1	15
aag tca tgt gcg aag aag cct aaa gag gag tcc cag aga gaa ctc aag	96
Lys Ser Cys Gly Lys Lys Pro Lys Glu Glu Ser Gln Arg Glu Leu Lys	
20	30
gga aag ata gac acc ctc acc ttg aag ctg aac gag aaa tcc aaa gag	144
Gly Lys Ile Asp Thr Leu Thr Leu Lys Leu Asn Glu Lys Ser Lys Glu	
35	45
cag gag gag ctt cta cag lys aag aat cag aac ctc caa gaa gcc ctg caa	192
Gln Glu Glu Leu Leu Gln Lys Asn Gln Asn Leu Gln Glu Ala Leu Gln	
50	60
aga gct gca aac ttt tca ggt cct tgt tca caa gac tgg ctt tgg cat	240
Arg Ala Ala Asn Phe Ser Gly Pro Cys Pro Gln Asp Trp Leu Trp His	
65	80
aaa gaa aac tgt tac ctc ttc cat ggg ccc ttt agc tgg gaa aaa aac	288
Lys Glu Asn Cys Tyr Leu Phe His Gly Pro Phe Ser Trp Glu Lys Asn	
85	95
cgg cag acc tgc caa tct ttg ggt ggc cag tta cta caa att aat ggt	336
Arg Gln Thr Cys Gln Ser Leu Gly Gly Gln Leu Leu Gln Ile Asn Gly	
100	110
gca gat gat ctg aca ttc atc tta caa gca att tcc cat acc acc tcc	384
Ala Asp Asp Leu Thr Phe Ile Leu Gln Ala Ile Ser His Thr Thr Ser	
115	125
cca ttc tgg att gga ttg cat cgg aag aag cct gcc caa cca tgg cta	432
Pro Phe Trp Ile Gly Leu His Arg Lys Lys Pro Gly Gln Pro Trp Leu	
130	140
tgg gag aat gga act cct ttg aat ttt caa ttc ttt aag acc agg ggc	480
Trp Glu Asn Gly Thr Pro Leu Asn Phe Gln Phe Phe Lys Thr Arg Gly	
145	160
gtt tct tta cag cta tat tca tca ggc aac tgt gca tac ctt caa gac	528
Val Ser Leu Gln Leu Tyr Ser Ser Gly Asn Cys Ala Tyr Leu Gln Asp	
165	175
gga gct gtg ttc gct gaa aac tgc att cta att gca ttc agc ata tgt	576
Gly Ala Val Phe Ala Glu Asn Cys Ile Leu Ile Ala Phe Ser Ile Cys	
180	190
cag aag aag aca aat cat ttg caa att tag	600
Gln Lys Lys Thr Asn His Leu Gln Ile	
195	200

Figure 11

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Isoform 9																
atg	act	ttt	gat	gac	aag	atg	aag	cct	ggc	aat	gac	gag	cct	gat	cag	48
Met	Thr	Phe	Asp	Asp	Lys	Met	Lys	Pro	Ala	Asn	Asp	Glu	Pro	Asp	Gln	
1				5					10					15		
aag	tca	tgt	ggc	aag	aag	cct	aaa	ggt	cct	tgt	cca	caa	gac	tgg	ctc	96
Lys	Ser	Cys	Gly	Lys	Lys	Pro	Lys	Gly	Pro	Cys	Pro	Gln	Asp	Trp	Leu	
			20					25					30			
tgg	cat	aaa	gaa	aac	tgt	tac	ctc	ttc	cat	ggg	ccc	ttt	agc	tgg	gaa	144
Trp	His	Lys	Glu	Asn	Cys	Tyr	Leu	Phe	His	Gly	Pro	Phe	Ser	Trp	Glu	
		35					40					45				
aaa	aac	egg	cag	acc	tgc	caa	tct	ttg	ggt	ggc	cag	tta	cta	caa	att	192
Lys	Asn	Arg	Gln	Thr	Cys	Gln	Ser	Leu	Gly	Gly	Gln	Leu	Leu	Gln	Ile	
	50					55					60					
aat	ggt	gca	gat	gat	ctg	aca	ttc	atc	tta	caa	gca	att	tcc	cat	acc	240
Asn	Gly	Ala	Asp	Asp	Leu	Thr	Phe	Ile	Leu	Gln	Ala	Ile	Ser	His	Thr	
65					70				75					80		
acc	tcc	cca	ttc	tgg	att	gga	ttg	cat	cgg	aag	aag	cct	ggc	caa	cca	288
Thr	Ser	Pro	Phe	Trp	Ile	Gly	Leu	His	Arg	Lys	Lys	Pro	Gly	Gln	Pro	
				85					90				95			
tgg	cta	tgg	gag	aat	gga	act	cct	ttg	aat	ttt	caa	ttc	ttt	aag	acc	336
Trp	Leu	Trp	Glu	Asn	Gly	Thr	Pro	Leu	Asn	Phe	Gln	Phe	Phe	Lys	Thr	
			100				105					110				
agg	ggc	gtt	tct	tta	cag	cta	tat	tca	tca	ggc	aac	tgt	gca	tac	ctt	384
Arg	Gly	Val	Ser	Leu	Gln	Leu	Tyr	Ser	Ser	Gly	Asn	Cys	Ala	Tyr	Leu	
		115					120				125					
caa	gac	gga	gct	gtg	ttc	gct	gaa	aac	tgc	att	cta	att	gca	ttc	agc	432
Gln	Asp	Gly	Ala	Val	Phe	Ala	Glu	Asn	Cys	Ile	Leu	Ile	Ala	Phe	Ser	
	130					135					140					
ata	tgt	cag	aag	aag	aca	aat	cat	ttg	caa	att	tag					480
Ile	Cys	Gln	Lys	Lys	Thr	Asn	His	Leu	Gln	Ile						
145					150				155							

Figure 12

A.

Isoform 1 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 1 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE
 Isoform 1 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 3 (R1) QSKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 3 (R3) ESQRELKGKIDTLTLKLNEKSKEQ...
 Isoform 4 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 7 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE
 Isoform 7 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 8 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG

B.

Isoform 1 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 3 (R1) QSKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 4 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANFSG

C.

Isoform 1 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE
 Isoform 7 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE

D.

Isoform 1 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 3 (R3) ESQRELKGKIDTLTLKLNEKSKEQ...
 Isoform 7 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 8 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG

E.

Isoform 1 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 1 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE
 Isoform 1 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 3 (R1) QSKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANSSE
 Isoform 3 (R3) ESQRELKGKIDTLTLKLNEKSKEQ...
 Isoform 4 (R1) ESKKELKGKIDTLTQKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 7 (R2) ESQRELKGKIDTITRKLDEKSKEQEELLQMIQNLQEALQRAANSSE
 Isoform 7 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 Isoform 8 (R3) ESQRELKGKIDTLTLKLNEKSKEQEELLQKNQNLQEALQRAANFSG
 human ESENELKEMIETLARKLNEKSKEQEMELHHQNLNLQETLKKRVANCSA

Figure 13

